

SEQUENCE LISTING

<110> BML, INC.

<120> Method of Detecting Risk Factor for Onset of Diabetes

<130> PBM37

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<170> PatentIn Ver. 2.0

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Met Ala Asn Cys Glu Phe Ser Pro Val Ser Gly Asp Lys Pro Cys Cys  
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cgg ctc tct agg aga gcc caa ctc tgt ctt ggc gtc agt atc ctg gtc 96  
Arg Leu Ser Arg Arg Ala Gln Leu Cys Leu Gly Val Ser Ile Leu Val  
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ctg atc ctc gtc gtg ctc gcg gtg gtc gtc ccg agg tgg cgc cag 144  
Leu Ile Leu Val Val Leu Ala Val Val Val Pro Arg Trp Arg Gln  
35 40 45

cag tgg agc ggt ccg ggc acc acc aag cgc ttt ccc gag acc gtc ctg 192  
Gln Trp Ser Gly Pro Gly Thr Thr Lys Arg Phe Pro Glu Thr Val Leu  
50 55 60

gcg cga tgc gtc aag tac act gaa att cat cct gag atg aga cat gta 240

Ala Arg Cys Val Lys Tyr Thr Glu Ile His Pro Glu Met Arg His Val			
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gac tgc caa agt gta tgg gat gct ttc aag ggt gca ttt att tca aaa			288
Asp Cys Gln Ser Val Trp Asp Ala Phe Lys Gly Ala Phe Ile Ser Lys			
85	90	95	
cat cct tgc aac att act gaa gaa gac tat cag cca cta atg aag ttg			336
His Pro Cys Asn Ile Thr Glu Glu Asp Tyr Gln Pro Leu Met Lys Leu			
100	105	110	
gga act cag acc gta cct tgc aac aag att ctt ctt tgg agc aga ata			384
Gly Thr Gln Thr Val Pro Cys Asn Lys Ile Leu Leu Trp Ser Arg Ile			
115	120	125	
aaa gat ctg gcc cat cag ttc aca cag gtc cag cggt gac atg ttc acc			432
Lys Asp Leu Ala His Gln Phe Thr Gln Val Gln Arg Asp Met Phe Thr			
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Leu Glu Asp Thr Leu Leu Gly Tyr Leu Ala Asp Asp Leu Thr Trp Cys			
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ggt gaa ttc aac act tcc aaa ata aac tat caa tct tgc cca gac tgg			528
Gly Glu Phe Asn Thr Ser Lys Ile Asn Tyr Gln Ser Cys Pro Asp Trp			
165	170	175	
aga aag gac tgc agc aac aac cct gtt tca gta ttc tgg aaa acg gtt			576
Arg Lys Asp Cys Ser Asn Asn Pro Val Ser Val Phe Trp Lys Thr Val			
180	185	190	
tcc cgc agg ttt gca gaa gct gcc tgt gat gtg gtc cat gtg atg ctc			624
Ser Arg Arg Phe Ala Glu Ala Ala Cys Asp Val Val His Val Met Leu			
195	200	205	
aat gga tcc cgc agt aaa atc ttt gac aaa aac agc act ttt ggg agt			672
Asn Gly Ser Arg Ser Lys Ile Phe Asp Lys Asn Ser Thr Phe Gly Ser			
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gtg gaa gtc cat aat ttg caa cca gag aag gtt cag aca cta gag gcc			720
Val Glu Val His Asn Leu Gln Pro Glu Lys Val Gln Thr Leu Glu Ala			
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tgg gtg ata cat ggt gga aga gaa gat tcc aga gac tta tgc cag gat			768
Trp Val Ile His Gly Gly Arg Glu Asp Ser Arg Asp Leu Cys Gln Asp			
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ccc acc ata aaa gag ctg gaa tcg att ata agc aaa agg aat att caa			816
Pro Thr Ile Lys Glu Leu Glu Ser Ile Ile Ser Lys Arg Asn Ile Gln			

260

265

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 Phe Ser Cys Lys Asn Ile Tyr Arg Pro Asp Lys Phe Leu Gln Cys Val  
 275 280 285

aaa aat cct gag gat tca tct tgc aca tct gag atc tgagccagtc 910  
 Lys Asn Pro Glu Asp Ser Ser Cys Thr Ser Glu Ile  
 290 295 300

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 35 40 45

Gln Trp Ser Gly Pro Gly Thr Thr Lys Arg Phe Pro Glu Thr Val Leu  
 50 55 60

Ala Arg Cys Val Lys Tyr Thr Glu Ile His Pro Glu Met Arg His Val  
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Asp Cys Gln Ser Val Trp Asp Ala Phe Lys Gly Ala Phe Ile Ser Lys  
 85 90 95

His Pro Cys Asn Ile Thr Glu Glu Asp Tyr Gln Pro Leu Met Lys Leu  
 100 105 110  
 Gly Thr Gln Thr Val Pro Cys Asn Lys Ile Leu Leu Trp Ser Arg Ile  
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 Lys Asp Leu Ala His Gln Phe Thr Gln Val Gln Arg Asp Met Phe Thr  
 130 135 140  
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 145 150 155 160  
 Gly Glu Phe Asn Thr Ser Lys Ile Asn Tyr Gln Ser Cys Pro Asp Trp  
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 Arg Lys Asp Cys Ser Asn Asn Pro Val Ser Val Phe Trp Lys Thr Val  
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 Ser Arg Arg Phe Ala Glu Ala Ala Cys Asp Val Val His Val Met Leu  
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 210 215 220  
 Val Glu Val His Asn Leu Gln Pro Glu Lys Val Gln Thr Leu Glu Ala  
 225 230 235 240  
 Trp Val Ile His Gly Gly Arg Glu Asp Ser Arg Asp Leu Cys Gln Asp  
 245 250 255  
 Pro Thr Ile Lys Glu Leu Glu Ser Ile Ile Ser Lys Arg Asn Ile Gln  
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